Reply to Stradner *et al.*: Equilibrium clusters are absent in concentrated lysozyme solutions

In their article (1), Stradner et al. stated, "Surprisingly, the peak position... is essentially independent" of concentration c referring to figure 1b in ref. 1 displaying six datasets from c = 36-273 mg/ml. In the main text of our article (2) a clear peak shift in the range of up to c = 200 mg/ml was demonstrated, and supporting information (SI) Fig. S9 (2) specifically presented systematic shifts for six concentrations (c =20-250 mg/ml). We thus disagree with the claim of "no significant differences" between the data presented in refs. 1 and 2. In their letter (3), Stradner et al. partially revoke their original statement (1) to say, "we both observe the usual shift in peak position at low concentrations" and argue about concentrated solutions without quantifying this notion. The peak shift is indeed less pronounced at higher concentrations corresponding to percolating clusters observed in the simulations (4), but this scenario is very different from the compact equilibrium clusters depicted in ref. 1. Our experimental data do not reveal typical small angle scattering features of equilibrium clusters as demonstrated in ref. 2. Instead the data are well described by physically meaningful dimensions and volume fraction of monomers, and this does allow one to reliably distinguish between monomers and clusters. In contrast, simulations of structure factors alone (1, 4) are incomplete because they do not take into account the form factor of the expected clusters but only that of the monomers. Summarizing, we observed no evidence for equilibrium clusters in concentrated lysozyme solutions for conditions similar to those reported in ref. 1. Formation of clusters at yet higher concentrations was neither studied nor questioned in our article (2).

Anuj Shukla^a, Efstratios Mylonas^b, Emanuela Di Cola^a, Stephanie Finet^a, Peter Timmins^c, Theyencheri Narayanan^a, and Dmitri I. Svergun^{b,d,1}

^aEuropean Synchrotron Radiation Facility, 6 Rue Jules Horowitz, F-38043, Grenoble Cedex 9, France; ^bEuropean Molecular Biology Laboratory, Hamburg Outstation, c/o DESY, Notkestrasse 85, D-22603 Hamburg, Germany; ^cInstitut Laue-Langevin, 6 Rue Jules Horowitz, F-38042, Grenoble Cedex 9, France; and ^dInstitute of Crystallography, Russian Academy of Sciences, Leninsky pr. 59, 117333 Moscow, Russia

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¹To whom correspondence should be addressed. E-mail: svergun@embl-hamburg.de. © 2008 by The National Academy of Sciences of the USA